SEQUENCE LISTING

(1) GENERAL INFORMATION:

- (i) APPLICANT: Murphy, Brian R. Collins, Peter L. Whitehead, Stephen S. Bukreyev, Alexander A. Juhasz, Katalin
- (ii) TITLE OF INVENTION: PRODUCTION OF ATTENUATED RESPIRATORY SYNCYTIAL VIRUS VACCINES FROM CLONED NUCLEOTIDE SEQUENCES
- (iii) NUMBER OF SEQUENCES: 14
- (iv) CORRESPONDENCE ADDRESS:
 - (A) ADDRESSEE: Townsend and Townsend and Crew LLP
 - (B) STREET: Two Embarcadero Center, 8th Floor
 - (C) CITY: San Francisco
 (D) STATE: CA

 - (E) COUNTRY: USA
 - (F) ZIP: 94111-3834
- (v) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE: Floppy disk
 - (B) COMPUTER: IBM PC compatible
 - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
 - (D) SOFTWARE: PatentIn Release #1.0, Version #1.25
- (vi) CURRENT APPLICATION DATA:
 - (A) APPLICATION NUMBER: US (B) FILING DATE: 15-JUL-1997

 - (C) CLASSIFICATION:
- (vii) PRIOR APPLICATION DATA:
 - (A) APPLICATION NUMBER: US 60/047,634
 - (B) FILING DATE: 23-MAY-1997
- (vii) PRIOR APPLICATION DATA:
 - (A) APPLICATION NUMBER: US 60/046,141
 - (B) FILING DATE: 09-MAY-1997
- (vii) PRIOR APPLICATION DATA:
 - (A) APPLICATION NUMBER: US 60/021,773
 - (B) FILING DATE: 15-JUL-1996
- (viii) ATTORNEY/AGENT INFORMATION:
 - (A) NAME: Parmelee, Steven W.
 - (B) REGISTRATION NUMBER: 31,990
 - (C) REFERENCE/DOCKET NUMBER: 17634-000510
 - (ix) TELECOMMUNICATION INFORMATION:
 - (A) TELEPHONE: 206-467-9600
 - (B) TELEFAX: 415-576-0300
- (2) INFORMATION FOR SEQ ID NO:1:
 - (i) SEOUENCE CHARACTERISTICS:
 - (A) LENGTH: 15223 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: cDNA

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

| ACGCGAAAAA | ATGCGTACAA | CAAACTTGCA | TAAACCAAAA | AAATGGGGCA | AATAAGAATT | 60 |
|------------|------------|------------|------------|------------|------------|------|
| TGATAAGTAC | CACTTAAATT | TAACTCCCTT | GGTTAGAGAT | GGGCAGCAAT | TCATTGAGTA | 120 |
| TGATAAAAGT | TAGATTACAA | AATTTGTTTG | ACAATGATGA | AGTAGCATTG | AATAAAATT | 180 |
| CATGCTATAC | TGATAAATTA | ATACATTTAA | CTAATGCTTT | GGCTAAGGCA | GTGATACATA | 240 |
| CAATCAAATT | GAATGGCATT | GTGTTTGTGC | ATGTTATTAC | AAGTAGTGAT | ATTTGCCCTA | 300 |
| ATAATAATAT | TGTAGTAAAA | TCCAATTTCA | CAACAATGCC | AGTACTACAA | AATGGAGGTT | 360 |
| ATATATGGGA | AATGATGGAA | TTAACACATT | GCTCTCAACC | TAATGGTCTA | CTAGATGACA | 420 |
| ATTGTGAAAT | TAAATTCTCC | AAAAAACTAA | GTGATTCAAC | AATGACCAAT | TATATGAATC | 480 |
| AATTATCTGA | ATTACTTGGA | TTTGATCTTA | ATCCATAAAT | TATAATTAAT | ATCAACTAGC | 540 |
| AAATCAATGT | CACTAACACC | ATTAGTTAAT | ATAAAACTTA | ACAGAAGACA | AAAATGGGGC | 600 |
| AAATAAATCA | ATTCAGCCAA | CCCAACCATG | GACACAACCC | ACAATGATAA | TACACCACAA | 660 |
| AGACTGATGA | TCACAGACAT | GAGACCGTTG | TCACTTGAGA | CCATAATAAC | ATCACTAACC | 720 |
| AGAGACATCA | TAACACACAA | ATTTATATAC | TTGATAAATC | ATGAATGCAT | AGTGAGAAAA | 780 |
| CTTGATGAAA | AGCAGGCCAC | ATTTACATTC | CTGGTCAACT | ATGAAATGAA | ACTATTACAC | 840 |
| AAAGTAGGAA | GCACTAAATA | TAAAAAATAT | ACTGAATACA | ACACAAAATA | TGGCACTTTC | 900 |
| CCTATGCCAA | TATTCATCAA | TCATGATGGG | TTCTTAGAAT | GCATTGGCAT | TAAGCCTACA | 960 |
| AAGCATACTC | CCATAATATA | CAAGTATGAT | CTCAATCCAT | AAATTTCAAC | ACAATATTCA | 1020 |
| САСААТСТАА | AACAACAACT | CTATGCATAA | CTATACTCCA | TAGTCCAGAT | GGAGCCTGAA | 1080 |
| AATTATAGTA | ATTTAAAACT | TAAGGAGAGA | TATAAGATAG | AAGATGGGGC | AAATACAACC | 1140 |
| ATGGCTCTTA | GCAAAGTCAA | GTTGAATGAT | ACACTCAACA | AAGATCAACT | TCTGTCATCC | 1200 |
| AGCAAATACA | CCATCCAACG | GAGCACAGGA | GATAGTATTG | ATACTCCTAA | TTATGATGTG | 1260 |
| CAGAAACACA | TCAATAAGTT | ATGTGGCATG | TTATTAATCA | CAGAAGATGC | TAATCATAAA | 1320 |
| TTCACTGGGT | TAATAGGTAT | GTTATATGCG | ATGTCTAGGT | TAGGAAGAGA | AGACACCATA | 1380 |
| AAAATACTCA | GAGATGCGGG | ATATCATGTA | AAAGCAAATG | GAGTAGATGT | AACAACACAT | 1440 |
| CGTCAAGACA | TTAATGGAAA | AGAAATGAAA | TTTGAAGTGT | TAACATTGGC | AAGCTTAACA | 1500 |
| ACTGAAATTC | AAATCAACAT | TGAGATAGAA | TCTAGAAAAT | CCTACAAAAA | AATGCTAAAA | 1560 |
| GAAATGGGAG | AGGTAGCTCC | AGAATACAGG | CATGACTCTC | CTGATTGTGG | GATGATAATA | 1620 |
| TTATGTATAG | CAGCATTAGT | AATAACTAAA | TTAGCAGCAG | GGGACAGATC | TGGTCTTACA | 1680 |
| GCCGTGATTA | GGAGAGCTAA | TAATGTCCTA | AAAAATGAAA | TGAAACGTTA | CAAAGGCTTA | 1740 |
| CTACCCAAGG | ACATAGCCAA | CAGCTTCTAT | GAAGTGTTTG | AAAAACATCC | CCACTTTATA | 1800 |
| GATGTTTTTG | TTCATTTTGG | TATAGCACAA | TCTTCTACCA | GAGGTGGCAG | TAGAGTTGAA | 1860 |
| GGGATTTTTG | CAGGATTGTT | TATGAATGCC | TATGGTGCAG | GGCAAGTGAT | GTTACGGTGG | 1920 |
| GGAGTCTTAG | CAAAATCAGT | TAAAAATATT | ATGTTAGGAC | ATGCTAGTGT | GCAAGCAGAA | 1980 |
| ATGGAACAAG | TTGTTGAGGT | TTATGAATAT | GCCCAAAAAT | TGGGTGGTGA | AGCAGGATTC | 2040 |

TACCATATAT TGAACAACCC AAAAGCATCA TTATTATCTT TGACTCAATT TCCTCACTTC 2100 TCCAGTGTAG TATTAGGCAA TGCTGCTGGC CTAGGCATAA TGGGAGAGTA CAGAGGTACA 2160 CCGAGGAATC AAGATCTATA TGATGCAGCA AAGGCATATG CTGAACAACT CAAAGAAAAT 2220 GGTGTGATTA ACTACAGTGT ACTAGACTTG ACAGCAGAAG AACTAGAGGC TATCAAACAT 2280 CAGCTTAATC CAAAAGATAA TGATGTAGAG CTTTGAGTTA ATAAAAAATG GGGCAAATAA 2340 ATCATCATGG AAAAGTTTGC TCCTGAATTC CATGGAGAAG ATGCAAACAA CAGGGCTACT 2400 AAATTCCTAG AATCAATAAA GGGCAAATTC ACATCACCCA AAGATCCCAA GAAAAAAGAT 2460 AGTATCATAT CTGTCAACTC AATAGATATA GAAGTAACCA AAGAAAGCCC TATAACATCA 2520 AATTCAACTA TTATCAACCC AACAAATGAG ACAGATGATA CTGCAGGGAA CAAGCCCAAT 2580 TATCAAAGAA AACCTCTAGT AAGTTTCAAA GAAGACCCTA CACCAAGTGA TAATCCCTTT 2640 TCTAAACTAT ACAAAGAAAC CATAGAAACA TTTGATAACA ATGAAGAAGA ATCCAGCTAT 2700 2760 TCATACGAAG AAATAAATGA TCAGACAAAC GATAATATAA CAGCAAGATT AGATAGGATT GATGAAAAAT TAAGTGAAAT ACTAGGAATG CTTCACACAT TAGTAGTGGC AAGTGCAGGA 2820 CCTACATCTG CTCGGGATGG TATAAGAGAT GCCATGGTTG GTTTAAGAGA AGAAATGATA 2880 GAAAAATCA GAACTGAAGC ATTAATGACC AATGACAGAT TAGAAGCTAT GGCAAGACTC 2940 AGGAATGAGG AAAGTGAAAA GATGGCAAAA GACACATCAG ATGAAGTGTC TCTCAATCCA 3000 ACATCAGAGA AATTGAACAA CCTATTGGAA GGGAATGATA GTGACAATGA TCTATCACTT 3060 GAAGATITCT GATTAGTTAC CAATCTTCAC ATCAACACAC AATACCAACA GAAGACCAAC 3120 3180 AAAACAACCA GCCAATCCAA AACTAACCAC CCGGAAAAAA TCTATAATAT AGTTACAAAA 3240 AAAGGAAAGG GTGGGGCAAA TATGGAAACA TACGTGAACA AGCTTCACGA AGGCTCCACA 3300 TACACAGCTG CTGTTCAATA CAATGTCTTA GAAAAAGACG ATGACCCTGC ATCACTTACA 3360 ATATGGGTGC CCATGTTCCA ATCATCTATG CCAGCAGATT TACTTATAAA AGAACTAGCT 3420 AATGTCAACA TACTAGTGAA ACAAATATCC ACACCCAAGG GACCTTCACT AAGAGTCATG 3480 ATAAACTCAA GAAGTGCAGT GCTAGCACAA ATGCCCAGCA AATTTACCAT ATGCGCTAAT 3540 GTGTCCTTGG ATGAAAGAAG CAAACTAGCA TATGATGTAA CCACACCCTG TGAAATCAAG 3600 GCATGTAGTC TAACATGCCT AAAATCAAAA AATATGTTGA CTACAGTTAA AGATCTCACT 3660 ATGAAGACAC TCAACCCTAC ACATGATATT ATTGCTTTAT GTGAATTTGA AAACATAGTA 3720 ACATCAAAAA AAGTCATAAT ACCAACATAC CTAAGATCCA TCAGTGTCAG AAATAAAGAT 3780 CTGAACACAC TTGAAAATAT AACAACCACT GAATTCAAAA ATGCTATCAC AAATGCAAAA 3840 ATCATCCCTT ACTCAGGATT ACTATTAGTC ATCACAGTGA CTGACAACAA AGGAGCATTC 3900 AAATACATAA AGCCACAAAG TCAATTCATA GTAGATCTTG GAGCTTACCT AGAAAAAGAA 3960 AGTATATATT ATGTTACCAC AAATTGGAAG CACACAGCTA CACGATTTGC AATCAAACCC 4020 ATGGAAGATT AACCTTTTTC CTCTACATCA GTGTGTTAAT TCATACAAAC TTTCTACCTA 4080 CATTCTTCAC TTCACCATCA CAATCACAAA CACTCTGTGG TTCAACCAAT CAAACAAAAC 4140 TTATCTGAAG TCCCAGATCA TCCCAAGTCA TTGTTTATCA GATCTAGTAC TCAAATAAGT 4200 TAATAAAAA TATACACATG GGGCAAATAA TCATTGGAGG AAATCCAACT AATCACAATA 4260 4320 TCTGTTAACA TAGACAAGTC CACACCCAT ACAGAATCAA CCAATGGAAA ATACATCCAT AACAATAGAA TTCTCAAGCA AATTCTGGCC TTACTTTACA CTAATACACA TGATCACAAC 4380 AATAATCTCT TIGCTAATCA TAATCTCCAT CATGATTGCA ATACTAAACA AACTTIGTGA 4440 ATATAACGTA TTCCATAACA AAACCTTTGA GTTACCAAGA GCTCGAGTCA ACACATAGCA 4500 TTCATCAATC CAACAGCCCA AAACAGTAAC CTTGCATTTA AAAATGAACA ACCCCTACCT 4560 CTTTACAACA CCTCATTAAC ATCCCACCAT GCAAACCACT ATCCATACTA TAAAGTAGTT 4620 AATTAAAAAT AGTCATAACA ATGAACTAGG ATATCAAGAC TAACAATAAC ATTGGGGCAA 4680 ATGCAAACAT GTCCAAAAAC AAGGACCAAC GCACCGCTAA GACATTAGAA AGGACCTGGG 4740 ACACTCTCAA TCATTTATTA TTCATATCAT CGTGCTTATA TAAGTTAAAT CTTAAATCTG 4800 TAGCACAAAT CACATTATCC ATTCTGGCAA TGATAATCTC AACTTCACTT ATAATTGCAG 4860 CCATCATATT CATAGCCTCG GCAAACCACA AAGTCACACC AACAACTGCA ATCATACAAG 4920 4980 ATGCAACAAG CCAGATCAAG AACACAACCC CAACATACCT CACCCAGAAT CCTCAGCTTG GAATCAGTCC CTCTAATCCG TCTGAAATTA CATCACAAAT CACCACCATA CTAGCTTCAA 5040 CAACACCAGG AGTCAAGTCA ACCCTGCAAT CCACAACAGT CAAGACCAAA AACACAACAA 5100 CAACTCAAAC ACAACCCAGC AAGCCCACCA CAAAACAACG CCAAAACAAA CCACCAAGCA 5160 AACCCAATAA TGATTTTCAC TTTGAAGTGT TCAACTTTGT ACCCTGCAGC ATATGCAGCA 5220 ACAATCCAAC CTGCTGGGCT ATCTGCAAAA GAATACCAAA CAAAAAACCA GGAAAGAAAA 5280 CCACTACCAA GCCCACAAAA AAACCAACCC TCAAGACAAC CAAAAAAGAT CCCAAACCTC 5340 AAACCACTAA ATCAAAGGAA GTACCCACCA CCAAGCCCAC AGAAGAGCCA ACCATCAACA 5400 CCACCAAAAC AAACATCATA ACTACACTAC TCACCTCCAA CACCACAGGA AATCCAGAAC 5460 TCACAAGTCA AATGGAAACC TTCCACTCAA CTTCCTCCGA AGGCAATCCA AGCCCTTCTC 5520 AAGTCTCTAC AACATCCGAG TACCCATCAC AACCTTCATC TCCACCCAAC ACACCACGCC 5580 AGTAGTTACT TAAAAACATA TTATCACAAA AGGCCTTGAC CAACTTAAAC AGAATCAAAA 5640 TAAACTCTGG GGCAAATAAC AATGGAGTTG CTAATCCTCA AAGCAAATGC AATTACCACA 5700 ATCCTCACTG CAGTCACATT TTGTTTTGCT TCTGGTCAAA ACATCACTGA AGAATTTTAT 5760 CAATCAACAT GCAGTGCAGT TAGCAAAGGC TATCTTAGTG CTCTGAGAAC TGGTTGGTAT 5820 ACCAGTGTTA TAACTATAGA ATTAAGTAAT ATCAAGAAAA ATAAGTGTAA TGGAACAGAT 5880 GCTAAGGTAA AATTGATAAA ACAAGAATTA GATAAATATA AAAATGCTGT AACAGAATTG 5940 CAGTTGCTCA TGCAAAGCAC ACAAGCAACA AACAATCGAG CCAGAAGAGA ACTACCAAGG 6000 TTTATGAATT ATACACTCAA CAATGCCAAA AAAACCAATG TAACATTAAG CAAGAAAAGG 6060 AAAAGAAGAT TTCTTGGTTT TTTGTTAGGT GTTGGATCTG CAATCGCCAG TGGCGTTGCT 6120 GTATCTAAGG TCCTGCACCT AGAAGGGGAA GTGAACAAGA TCAAAAGTGC TCTACTATCC 6180 ACAAACAAGG CTGTAGTCAG CTTATCAAAT GGAGTTAGTG TTTTAACCAG CAAAGTGTTA 6240 GACCTCAAAA ACTATATAGA TAAACAATTG TTACCTATTG TGAACAAGCA AAGCTGCAGC 6300 ATATCAAATA TAGAAACTGT GATAGAGTTC CAACAAAAGA ACAACAGACT ACTAGAGATT 6360 ACCAGGGAAT TTAGTGTTAA TGCAGGCGTA ACTACACCTG TAAGCACTTA CATGTTAACT 6420 AATAGTGAAT TATTGTCATT AATCAATGAT ATGCCTATAA CAAATGATCA GAAAAAGTTA 6480 ATGTCCAACA ATGTTCAAAT AGTTAGACAG CAAAGTTACT CTATCATGTC CATAATAAAA 6540 GAGGAAGTCT TAGCATATGT AGTACAATTA CCACTATATG GTGTTATAGA TACACCCTGT 6600 TGGAAACTAC ACACATCCCC TCTATGTACA ACCAACACAA AAGAAGGGTC CAACATCTGT 6660 TTAACAAGAA CTGACAGAGG ATGGTACTGT GACAATGCAG GATCAGTATC TTTCTTCCCA 6720 CAAGCTGAAA CATGTAAAGT TCAATCAAAT CGAGTATTTT GTGACACAAT GAACAGTTTA 6780 ACATTACCAA GTGAAGTAAA TCTCTGCAAT GTTGACATAT TCAACCCCAA ATATGATTGT 6840 AAAATTATGA CTTCAAAAAC AGATGTAAGC AGCTCCGTTA TCACATCTCT AGGAGCCATT 6900 6960 GTGTCATGCT ATGGCAAAAC TAAATGTACA GCATCCAATA AAAATCGTGG AATCATAAAG ACATTTTCTA ACGGGTGCGA TTATGTATCA AATAAAGGGG TGGACACTGT GTCTGTAGGT 7020 AACACATTAT ATTATGTAAA TAAGCAAGAA GGTAAAAGTC TCTATGTAAA AGGTGAACCA 7080 ATAATAAATT TCTATGACCC ATTAGTATTC CCCTCTGATG AATTTGATGC ATCAATATCT 7140 CAAGTCAACG AGAAGATTAA CCAGAGCCTA GCATTTATTC GTAAATCCGA TGAATTATTA 7200 CATAATGTAA ATGCTGGTAA ATCCACCACA AATATCATGA TAACTACTAT AATTATAGTG 7260 ATTATAGTAA TATTGTTATC ATTAATTGCT GTTGGACTGC TCTTATACTG TAAGGCCAGA 7320 AGCACACCAG TCACACTAAG CAAAGATCAA CTGAGTGGTA TAAATAATAT TGCATTTAGT 7380 AACTAAATAA AAATAGCACC TAATCATGTT CTTACAATGG TTTACTATCT GCTCATAGAC 7440 AACCCATCTG TCATTGGATT TTCTTAAAAT CTGAACTTCA TCGAAACTCT CATCTATAAA 7500 CCATCTCACT TACACTATTT AAGTAGATTC CTAGTTTATA GTTATATAAA ACACAATTGC 7560 ATGCCAGATT AACTTACCAT CTGTAAAAAT GAAAACTGGG GCAAATATGT CACGAAGGAA 7620 TCCTTGCAAA TTTGAAATTC GAGGTCATTG CTTAAATGGT AAGAGGTGTC ATTTTAGTCA 7680 TAATTATTT GAATGGCCAC CCCATGCACT GCTTGTAAGA CAAAACTTTA TGTTAAACAG 7740 7800 AATACTTAAG TCTATGGATA AAAGTATAGA TACCTTATCA GAAATAAGTG GAGCTGCAGA GTTGGACAGA ACAGAAGAGT ATGCTCTTGG TGTAGTTGGA GTGCTAGAGA GTTATATAGG 7860 ATCAATAAAC AATATAACTA AACAATCAGC ATGTGTTGCC ATGAGCAAAC TCCTCACTGA 7920 ACTCAATAGT GATGATATCA AAAAGCTGAG GGACAATGAA GAGCTAAATT CACCCAAGAT 7980 AAGAGTGTAC AATACTGTCA TATCATATAT TGAAAGCAAC AGGAAAAACA ATAAACAAAC 8040 TATCCATCTG TTAAAAAGAT TGCCAGCAGA CGTATTGAAG AAAACCATCA AAAACACATT 8100 GGATATCCAT AAGAGCATAA CCATCAACAA CCCAAAAGAA TCAACTGTTA GTGATACAAA 8160 TGACCATGCC AAAAATAATG ATACTACCTG ACAAATATCC TTGTAGTATA ACTTCCATAC 8220 TAATAACAAG TAGATGTAGA GTTACTATGT ATAATCAAAA GAACACACTA TATTTCAATC 8280 ANACCACCC ANATANCCAT ATGTACTCAC CGAATCANAC ATTCAATGAN ATCCATTGGA 8340 CCTCTCAAGA ATTGATTGAC ACAATTCAAA ATTTTCTACA ACATCTAGGT ATTATTGAGG 8400 ATATATATA AATATATATA TTAGTGTCAT AACACTCAAT TCTAACACTC ACCACATCGT 8460 TACATTATTA ATTCAAACAA TTCAAGTTGT GGGACAAAAT GGATCCCATT ATTAATGGAA 8520 ATTCTGCTAA TGTTTATCTA ACCGATAGTT ATTTAAAAGG TGTTATCTCT TTCTCAGAGT 8580 GTAATGCTTT AGGAAGTTAC ATATTCAATG GTCCTTATCT CAAAAATGAT TATACCAACT 8640 TAATTAGTAG ACAAAATCCA TTAATAGAAC ACATGAATCT AAAGAAACTA AATATAACAC 8700 AGTCCTTAAT ATCTAAGTAT CATAAAGGTG AAATAAAATT AGAAGAACCT ACTTATTTTC 8760 AGTCATTACT TATGACATAC AAGAGTATGA CCTCGTCAGA ACAGATTGCT ACCACTAATT 8820 TACTTAAAAA GATAATAAGA AGAGCTATAG AAATAAGTGA TGTCAAAGTC TATGCTATAT 8880 TGAATAAACT AGGGCTTAAA GAAAAGGACA AGATTAAATC CAACAATGGA CAAGATGAAG 8940 ACAACTCAGT TATTACGACC ATAATCAAAG ATGATATACT TTCAGCTGTT AAAGATAATC 9000 AATCTCATCT TAAAGCAGAC AAAAATCACT CTACAAAACA AAAAGACACA ATCAAAACAA 9060 CACTCTTGAA GAAATTGATG TGTTCAATGC AACATCCTCC ATCATGGTTA ATACATTGGT 9120 TTAACTTATA CACAAAATTA AACAACATAT TAACACAGTA TCGATCAAAT GAGGTAAAAA 9180 ACCATGGGTT TACATTGATA GATAATCAAA CTCTTAGTGG ATTTCAATTT ATTTTGAACC 9240 AATATGGTTG TATAGTTTAT CATAAGGAAC TCAAAAGAAT TACTGTGACA ACCTATAATC 9300 AATTCTTGAC ATGGAAAGAT ATTAGCCTTA GTAGATTAAA TGTTTGTTTA ATTACATGGA 9360 TTAGTAACTG CTTGAACACA TTAAATAAAA GCTTAGGCTT AAGATGCGGA TTCAATAATG 9420 TTATCTTGAC ACAACTATTC CTTTATGGAG ATTGTATACT AAAGCTATTT CACAATGAGG 9480 GGTTCTACAT AATAAAAGAG GTAGAGGGAT TTATTATGTC TCTAATTTTA AATATAACAG 9540 9600 AAGAAGATCA ATTCAGAAAA CGATTTTATA ATAGTATGCT CAACAACATC ACAGATGCTG CTAATAAGC TCAGAAAAAT CTGCTATCAA GAGTATGTCA TACATTATTA GATAAGACAG 9660 TGTCCGATAA TATAATAAAT GGCAGATGGA TAATTCTATT AAGTAAGTTC CTTAAATTAA 9720 9780 TTAAGCTTGC AGGTGACAAT AACCTTAACA ATCTGAGTGA ACTATATTTT TTGTTCAGAA TATTTGGACA CCCAATGGTA GATGAAAGAC AAGCCATGGA TGCTGTTAAA ATTAATTGCA 9840 ATGAGACCAA ATTTTACTTG TTAAGCAGTC TGAGTATGTT AAGAGGTGCC TTTATATATA 9900 GAATTATAAA AGGGTTTGTA AATAATTACA ACAGATGGCC TACTTTAAGA AATGCTATTG 9960 TTTTACCCTT AAGATGGTTA ACTTACTATA AACTAAACAC TTATCCTTCT TTGTTGGAAC 10020 10080 TTACAGAAAG AGATTTGATT GTGTTATCAG GACTACGTTT CTATCGTGAG TTTCGGTTGC CTAAAAAGT GGATCTTGAA ATGATTATAA ATGATAAAGC TATATCACCT CCTAAAAATT 10140 TGATATGGAC TAGTTTCCCT AGAAATTACA TGCCATCACA CATACAAAAC TATATAGAAC 10200 ATGAAAAATT AAAATTTTCC GAGAGTGATA AATCAAGAAG AGTATTAGAG TATTATTTAA 10260 GAGATAACAA ATTCAATGAA TGTGATTTAT ACAACTGTGT AGTTAATCAA AGTTATCTCA 10320 ACAACCTAA TCATGTGGTA TCATTGACAG GCAAAGAAAG AGAACTCAGT GTAGGTAGAA 10380 TGTTTGCAAT GCAACCGGGA ATGTTCAGAC AGGTTCAAAT ATTGGCAGAG AAAATGATAG 10440 CTGAAAACAT TTTACAATTC TTTCCTGAAA GTCTTACAAG ATATGGTGAT CTAGAACTAC 10500 AAAAATATT AGAACTGAAA GCAGGAATAA GTAACAAATC AAATCGCTAC AATGATAATT 10560 ACAACAATTA CATTAGTAAG TGCTCTATCA TCACAGATCT CAGCAAATTC AATCAAGCAT 10620 TTCGATATGA AACGTCATGT ATTTGTAGTG ATGTGCTGGA TGAACTGCAT GGTGTACAAT 10680 CTCTATTTTC CTGGTTACAT TTAACTATTC CTCATGTCAC AATAATATGC ACATATAGGC 10740 ATGCACCCC CTATATAGGA GATCATATTG TAGATCTTAA CAATGTAGAT GAACAAAGTG 10800 GATTATATAG ATATCACATG GGTGGCATCG AAGGGTGGTG TCAAAAACTA TGGACCATAG 10860 AAGCTATATC ACTATTGGAT CTAATATCTC TCAAAGGGAA ATTCTCAATT ACTGCTTTAA 10920 TTAATGGTGA CAATCAATCA ATAGATATAA GCAAACCAAT CAGACTCATG GAAGGTCAAA 10980 CTCATGCTCA AGCAGATTAT TTGCTAGCAT TAAATAGCCT TAAATTACTG TATAAAGAGT 11040 ATGCAGGCAT AGGCCACAAA TTAAAAGGAA CTGAGACTTA TATATCACGA GATATGCAAT 11100 11160 TTATGAGTAA AACAATTCAA CATAACGGTG TATATTACCC AGCTAGTATA AAGAAAGTCC TAAGAGTGGG ACCGTGGATA AACACTATAC TTGATGATTT CAAAGTGAGT CTAGAATCTA 11220 TAGGTAGTTT GACACAAGAA TTAGAATATA GAGGTGAAAG TCTATTATGC AGTTTAATAT 11280 TTAGAAATGT ATGGTTATAT AATCAGATTG CTCTACAATT AAAAAATCAT GCATTATGTA 11340 ACAATAAACT ATATTTGGAC ATATTAAAGG TTCTGAAACA CTTAAAAACC TTTTTTAATC 11400 TTGATAATAT TGATACAGCA TTAACATTGT ATATGAATTT ACCCATGTTA TTTGGTGGTG 11460 GTGATCCCAA CTTGTTATAT CGAAGTTTCT ATAGAAGAAC TCCTGACTTC CTCACAGAGG 11520 CTATAGTTCA CTCTGTGTTC ATACTTAGTT ATTATACAAA CCATGACTTA AAAGATAAAC 11580 TTCAAGATCT GTCAGATGAT AGATTGAATA AGTTCTTAAC ATGCATAATC ACGTTTGACA 11640 AAAACCCTAA TGCTGAATTC GTAACATTGA TGAGAGATCC TCAAGCTTTA GGGTCTGAGA 11700 GACAAGCTAA AATTACTAGC GAAATCAATA GACTGGCAGT TACAGAGGTT TTGAGTACAG 11760 . CTCCAAACAA AATATTCTCC AAAAGTGCAC AACATTATAC TACTACAGAG ATAGATCTAA 11820 ATGATATTAT GCAAAATATA GAACCTACAT ATCCTCATGG GCTAAGAGTT GTTTATGAAA 11880 GTTTACCCTT TTATAAAGCA GAGAAAATAG TAAATCTTAT ATCAGGTACA AAATCTATAA 11940 CTAACATACT GGAAAAAACT TCTGCCATAG ACTTAACAGA TATTGATAGA GCCACTGAGA 12000 TGATGAGGAA AAACATAACT TTGCTTATAA GGATACTTCC ATTGGATTGT AACAGAGATA 12060 AAAGAGAGAT ATTGAGTATG GAAAACCTAA GTATTACTGA ATTAAGCAAA TATGTTAGGG 12120 AAAGATCTTG GTCTTTATCC AATATAGTTG GTGTTACATC ACCCAGTATC ATGTATACAA 12180 TGGACATCAA ATATACTACA AGCACTATAT CTAGTGGCAT AATTATAGAG AAATATAATG 12240 TTAACAGTTT AACACGTGGT GAGAGAGGAC CCACTAAACC ATGGGTTGGT TCATCTACAC 12300 AAGAGAAAA AACAATGCCA GTTTATAATA GACAAGTCTT AACCAAAAAA CAGAGAGATC 12360 AAATAGATCT ATTAGCAAAA TTGGATTGGG TGTATGCATC TATAGATAAC AAGGATGAAT 12420 TCATGGAAGA ACTCAGCATA GGAACCCTTG GGTTAACATA TGAAAAGGCC AAGAAATTAT 12480 TTCCACATA TTTAAGTGTC AATTATTTGC ATCGCCTTAC AGTCAGTAGT AGACCATGTG 12540

AATTCCCTGC ATCAATACCA GCTTATAGAA CAACAAATTA TCACTTTGAC ACTAGCCCTA 12600 TTAATCGCAT ATTAACAGAA AAGTATGGTG ATGAAGATAT TGACATAGTA TTCCAAAACT 12660 GTATAAGCTT TGGCCTTAGT TTAATGTCAG TAGTAGAACA ATTTACTAAT GTATGTCCTA 12720 ACAGAATTAT TCTCATACCT AAGCTTAATG AGATACATTT GATGAAACCT CCCATATTCA 12780 CAGGTGATGT TGATATTCAC AAGTTAAAAC AAGTGATACA AAAACAGCAT ATGTTTTTAC 12840 CAGACAAAAT AAGTTTGACT CAATATGTGG AATTATTCTT AAGTAATAAA ACACTCAAAT 12900 CTGGATCTCA TGTTAATTCT AATTTAATAT TGGCACATAA AATATCTGAC TATTTTCATA 12960 ATACTTACAT TTTAAGTACT AATTTAGCTG GACATTGGAT TCTGATTATA CAACTTATGA 13020 AAGATTCTAA AGGTATTTTT GAAAAAGATT GGGGAGAGGG ATATATAACT GATCATATGT 13080 TTATTAATTT GAAAGTTITC TTCAATGCTT ATAAGACCTA TCTCTTGTGT TTTCATAAAG 13140 GTTATGGCAA AGCAAAGCTG GAGTGTGATA TGAACACTTC AGATCTTCTA TGTGTATTGG 13200 AATTAATAGA CAGTAGTTAT TGGAAGTCTA TGTCTAAGGT ATTTTTAGAA CAAAAAGTTA 13260 TCAAATACAT TCTTAGCCAA GATGCAAGTT TACATAGAGT AAAAGGATGT CATAGCTTCA 13320 AATTATGGTT TCTTAAACGT CTTAATGTAG CAGAATTCAC AGTTTGCCCT TGGGTTGTTA 13380 ACATAGATTA TCATCCAACA CATATGAAAG CAATATTAAC TTATATAGAT CTTGTTAGAA 13440 TGGGATTGAT AAATATAGAT AGAATACACA TTAAAAATAA ACACAAATTC AATGATGAAT 13500 TTTATACTTC TAATCTCTTC TACATTAATT ATAACTTCTC AGATAATACT CATCTATTAA 13560 CTAAACATAT AAGGATTGCT AATTCTGAAT TAGAAAATAA TTACAACAAA TTATATCATC 13620 CTACACCAGA AACCCTAGAG AATATACTAG CCAATCCGAT TAAAAGTAAT GACAAAAAGA 13680 CACTGAATGA CTATTGTATA GGTAAAAATG TTGACTCAAT AATGTTACCA TTGTTATCTA 13740 ATAAGAAGCT TATTAAATCG TCTGCAATGA TTAGAACCAA TTACAGCAAA CAAGATTTGT 13800 ATAATTTATT CCCTATGGTT GTGATTGATA GAATTATAGA TCATTCAGGC AATACAGCCA 13860 AATCCAACCA ACTITACACT ACTACTTCCC ACCAAATATC CTTAGTGCAC AATAGCACAT 13920 CACTTTACTG CATGCTTCCT TGGCATCATA TTAATAGATT CAATTTTGTA TTTAGTTCTA 13980 CAGGTTGTAA AATTAGTATA GAGTATATTT TAAAAGATCT TAAAATTAAA GATCCCAATT 14040 GTATAGCATT CATAGGTGAA GGAGCAGGGA ATTTATTATT GCGTACAGTA GTGGAACTTC 14100 ATCCTGACAT AAGATATATT TACAGAAGTC TGAAAGATTG CAATGATCAT AGTTTACCTA 14160 TTGAGTTTTT AAGGCTGTAC AATGGACATA TCAACATTGA TTATGGTGAA AATTTGACCA 14220 TTCCTGCTAC AGATGCAACC AACAACATTC ATTGGTCTTA TTTACATATA AAGTTTGCTG 14280 AACCTATCAG TCTTTTTGTC TGTGATGCCG AATTGTCTGT AACAGTCAAC TGGAGTAAAA 14340 TTATAATAGA ATGGAGCAAG CATGTAAGAA AGTGCAAGTA CTGTTCCTCA GTTAATAAAT 14400 GTATGTTAAT AGTAAAATAT CATGCTCAAG ATGATATTGA TITCAAATTA GACAATATAA 14460 CTATATTAAA AACTTATGTA TGCTTAGGCA GTAAGTTAAA GGGATCGGAG GTTTACTTAG 14520 TCCTTACAAT AGGTCCTGCG AATATATTCC CAGTATTTAA TGTAGTACAA AATGCTAAAT 14580 TGATACTATC AAGAACCAAA AATTTCATCA TGCCTAAGAA AGCTGATAAA GAGTCTATTG 14640 ATGCAAATAT TAAAAGTTTG ATACCCTTTC TTTGTTACCC TATAACAAAA AAAGGAATTA 14700 ATACTGCATT GTCAAAACTA AAGAGTGTTG TTAGTGGAGA TATACTATCA TATTCTATAG 14760 CTGGACGTAA TGAAGTTTTC AGCAATAAAC TTATAAATCA TAAGCATATG AACATCTTAA 14820 AATGGTTCAA TCATGTTTTA AATTTCAGAT CAACAGAACT AAACTATAAC CATTTATATA 14880 TGGTAGAATC TACATATCCT TACCTAAGTG AATTGTTAAA CAGCTTGACA ACCAATGAAC 14940 TTAAAAAACT GATTAAAATC ACAGGTAGTC TGTTATACAA CTTTCATAAT GAATAATGAA 15000 TAAAGATCTT ATAATAAAA TTCCCATAGC TATACACTAA CACTGTATTC AATTATAGTT 15060 ATTAAAAATT AAAAATCATA TAATTITTTA AATAACTITT AGTGAACTAA TCCTAAAGTT 15120 ATCATTTAA TCTTGGAGGA ATAAATTTAA ACCCTAATCT AATTGGTTTA TATGTGTATT 15180 15223 AACTAAATTA CGAGATATTA GTTTTTGACA CTTTTTTCT CGT

(2) INFORMATION FOR SEQ ID NO:2:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 15225 base pairs
- (B) TYPE: nucleic acid (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

| 60 | AATAAGAATT | AAATGGGGCA | CATTCGGAAA | CAAACTTGCA | ATGCGTACTA | ACGCGAAAAA |
|------|---------------------------|------------|------------|------------|------------|------------|
| 120 | TCACTGAGCA | GGGGTGCAAT | AATCAGAAAT | TAACCTTTTC | TATTTAAGTC | TGATAAGTGC |
| 180 | $\mathbf{TTAAAAATTA}^{'}$ | AGTAGCATTG | ACAATGACGA | AATTTATTTG | TAGATTACAA | TGATAAAGGT |
| 240 | GCAATACATA | AGCCAAAGCA | CCAATGCATT | ATTCTTCTGA | TGACAAATTA | CATGTTATAC |
| 300 | GTGTGCCCTG | AAGCAGTGAA | ATGTTATAAC | GTTTTTATAC | AAACGGTATA | CAATTAAATT |
| 360 | AACGGAGGAT | AATATTACAA | CAACAATGCC | TCTAACTTTA | TGTAGTAAAA | ATAACAACAT |
| 420 | ATGGATGATA | AAACGGTCTA | GCTCTCAATT | TTGACACACT | ATTGATTGAG | ACATATGGGA |
| 480 | TATATGAATC | AATGACTAAT | GTGACTCAGT | AAAAGACTAA | CAAATTTTCT | ATTGTGAAAT |
| 540 | CTAACTCAAT | TATGTTTAGT | ATTCATGAAT | CTTGATCTCA | TTTACTTGGG | AAATATCTGA |
| 600 | AATGGGGCAA | ATCAAAGGGA | АТАААААСТС | TTTAGTTAAT | TTATTACCAT | AGACATGTGT |
| 660 | CTATGCAAAG | GACAACACTA | CACTACAAAT | AAACTATGAG | CTAATCAATC | ATAAACTCAC |
| 720 | CTCTCACCAA | ATAATAACAT | GATGGATTCA | GACCCCTGTC | ACGGACATGA | ATTAATGATC |
| 780 | TAAGAAAACT | GAATGTATTG | GATAAACAAT | TCATATACTT | ACACACAAAT | AGAAATCATC |
| 840 | TACTGCACAA | GAGATGAAGC | AGTCAATTAT | TTACATTCTT | CAAGCTACAT | TGATGAAAGA |
| 900 | GCACTTTCCC | ACAAAATATG | TGAATATAAT | AGAAATACAC | ACCAAATACA | AGTAGGGAGT |
| 960 | AGCCTACAAA | ATTGGCATTA | TCTAGAATGT | ATGGCGGGTT | TTTATCAATC | CATGCCTATA |
| 1020 | ААААААССАА | ATTCCAACAA | CAACCCGTAA | AATATGACCT | ATAATATACA | ACACACTCCT |
| 1080 | GGAGCTAATC | TAGTTAAGAA | CAATGCTCAA | CCTCAAACAA | CCAAGCTATT | CCCAACCAAA |

| CGTTTTAGTA | ATTAAAAATA | AAAGTAAAGC | CAATAACATA | AATTGGGGCA | AATACAAAGA | 1140 |
|------------|--------------|-------------|--------------|--------------|--------------|------|
| TGGCTCTTAG | CAAAGTCAAG | TTAAATGATA | CATTAAATAA | GGATCAGCTG | CTGTCATCCA | 1200 |
| GCAAATACAC | TATTCAACGT | AGTACAGGAG | ATAATATTGA | CACTCCCAAT | TATGATGTGC | 1260 |
| AAAAACACCT | AAACAAACTA | TGTGGTATGC | TATTAATCAC | TGAAGATGCA | AATCATAAAT | 1320 |
| TCACAGGATT | AATAGGTATG | TTATATGCTA | TGTCCAGGTT | AGGAAGGGAA | GACACTATAA | 1380 |
| AGATACTTAA | AGATGCTGGA | TATCATGTTA | AAGCTAATGG | AGTAGATATA | ACAACATATC | 1440 |
| GTCAAGATAT | AAATGGAAAG | GAAATGAAAT | TCGAAGTATT | AACATTATCA | AGCTTGACAT | 1500 |
| | | | CTAGAAAATC | • | | 1560 |
| | | | 'ATGATTCTCC | | | 1620 |
| | | | TAGCAGCAGG | | | 1680 |
| | | | AAAATGAAAT | | | 1740 |
| | | | AAGTGTTTGA | | | 1800 |
| | | | CATCAACAAG | | | 1860 |
| | | | ATGGTTCAGG | | | 1920 |
| GAGTTTTAGC | CAAATCTGTA | AAAAATATCA | TGCTAGGTCA | TGCTAGTGTC | CAGGCAGAAA | 1980 |
| TGGAGCAAGI | TGTGGAAGTC | TATGAGTATG | CACAGAAGTT | GGGAGGAGAA | GCTGGATTCT | 2040 |
| ACCATATATI | GAACAATCCA | AAAGCATCAT | TGCTGTCATT | AACTCAATTI | CCTAACTTCT | 2100 |
| CAAGTGTGGT | CCTAGGCAAT | GCAGCAGGTC | TAGGCATAAT | GGGAGAGTAT | AGAGGTACGC | 2160 |
| CAAGAAACCA | GGATCTTTAT | GATGCAGCCA | AAGCATATGC | AGAGCAACTO | AAAGAAAATG | 2220 |
| GAGTAATAA | CTACAGTGTA | TTAGACTTA | CAGCAGAAGA | ATTGGAAGCC | ATAAAGAATC | 2280 |
| AACTCAACCC | TAAAGAAGAT | GATGTAGAGO | TTTAAGTTAA | CAAAAAATAC | GGGGCAAATA | 2340 |
| | | | | | ACAAAGCTAC | 2400 |
| CAAATTCCT | A GAATCAATAA | AGGGCAAGTT | CGCATCATCC | AAAGATCCT | AGAAGAAAGA | 2460 |
| TAGCATAATA | A TCTGTTAACT | CAATAGATAT | AGAAGTAACO | : AAAGAGAGCO | CGATAACATC | 2520 |
| TGGCACCAA | ATCATCAATC | CAACAAGTG | A AGCCGACAGT | ACCCCAGAA | A CCAAAGCCAA | 2580 |
| | | | | | ACAACCCTTT | 2640 |
| | | | | | AATCTAGCTA | 2700 |
| | | | | | C TAGATAGAAT | 2760 |
| | | | | | G CAAGTGCAGG | 2820 |
| ACCCACTTC | A GCTCGCGAT | GAATAAGAG | A TGCTATGGT | r ggtctgaga | G AAGAAATGAT | 2880 |
| | | | | | A TGGCAAGACT | 2940 |
| | | | | | C CTCTTAATCC | 3000 |
| | | | | | G ATCTGTCACT | 3060 |
| | | | | | T AAAACAGACA | 3120 |
| TCAATCCAT | T GAATCAACT | G CCAGACCGA | A CAAACAAAT | G TCCGTCAGC | G GAACCACCAA | 3180 |

CCAATCAATC AACCAACTGA TCCATCAGCA ACCTGACGAA ATTAACAATA TAGTAACAAA 3240 AAAAGAACAA GATGGGGCAA ATATGGAAAC ATACGTGAAC AAGCTTCACG AAGGCTCCAC 3300 ATACACAGCA GCTGTTCAGT ACAATGTTCT AGAAAAAGAT GATGATCCTG CATCACTAAC 3360 AATATGGGTG CCTATGTTCC AGTCATCTGT ACCAGCAGAC TTGCTCATAA AAGAACTTGC 3420 AAGCATCAAC ATACTAGTGA AGCAGATCTC TACGCCCAAA GGACCTTCAC TACGAGTCAC 3480 GATTAACTCA AGAAGTGCTG TGCTGGCTCA AATGCCTAGT AATTTCATCA TAAGCGCAAA 3540 TGTATCATTA GATGAAAGAA GCAAATTAGC ATATGATGTA ACTACACCTT GTGAAATCAA 3600 AGCATGCAGT CTAACATGCT TAAAAGTGAA AAGTATGTTA ACTACAGTCA AAGATCTTAC 3660 CATGAAGACA TTCAACCCCA CTCATGAGAT CATTGCTCTA TGTGAATTTG AAAATATTAT 3720 GACATCAAAA AGAGTAATAA TACCAACCTA TCTAAGACCA ATTAGTGTCA AAAACAAGGA 3780 TCTGAACTCA CTAGAAAACA TAGCAACCAC CGAATTCAAA AATGCTATCA CCAATGCGAA 3840 AATTATTCCC TATGCTGGAT TAGTATTAGT TATCACAGTT ACTGACAATA AAGGAGCATT 3900 CAAATATATC AAGCCACAGA GTCAATTTAT AGTAGATCTT GGTGCCTACC TAGAAAAAGA 3960 GAGCATATAT TATGTGACTA CTAATTGGAA GCATACAGCT ACACGTTTTT CAATCAAACC 4020 ACTAGAGGAT TAAATTTAAT TATCAACACT GAATGACAGG TCCACATATA TCCTCAAACT 4080 ACACACTATA TCCAAACATC ATGAACATCT ACACTACACA CTTCATCACA CAAACCAATC 4140 CCACTCAAAA TCCAAAATCA CTACCAGCCA CTATCTGCTA GACCTAGAGT GCGAATAGGT 4200 AAATAAAACC AAAATATGGG GTAAATAGAC ATTAGTTAGA GTTCAATCAA TCTCAACAAC 4260 CATTTATACC GCCAATTCAA TACATATACT ATAAATCTTA AAATGGGAAA TACATCCATC 4320 ACAATAGAAT TCACAAGCAA ATTTTGGCCC TATTTTACAC TAATACATAT GATCTTAACT 4380 CTAATCTCTT TACTAATTAT AATCACTATT ATGATTGCAA TACTAAATAA GCTAAGTGAA 4440 CATAAAACAT TCTGTAACAA TACTCTTGAA CTAGGACAGA TGCATCAAAT CAACACATAG 4500 TGCTCTACCA TCATGCTGTG TCAAATTATA ATCCTGTATA TATAAACAAA CAAATCCAAT 4560 CTTCTCACAG AGTCATGGTG TCGCAAAACC ACGCCAACTA TCATGGTAGC ATAGAGTAGT 4620 TATTTAAAAA TTAACATAAT GATGAATTAT TAGTATGGGA TCAAAAACAA CATTGGGGCA 4680 AATGCAACCA TGTCCAAACA CAAGAATCAA CGCACTGCCA GGACTCTAGA AAAGACCTGG 4740 GATACTCTCA ATCATCTAAT TGTAATATCC TCTTGTTTAT ACAGATTAAA TTTAAAATCT 4800 ATAGCACAAA TAGCACTATC AGTTCTGGCA ATGATAATCT CAACCTCTCT CATAATTGCA 4860 GCCATAATAT TCATCATCTC TGCCAATCAC AAAGTTACAC TAACAACGGT CACAGTTCAA 4920 ACAATAAAAA ACCACACTGA AAAAAACATC ACCACCTACC TTACTCAAGT CCCACCAGAA 4980 AGGGTTAGCT CATCCAAACA ACCTACAACC ACATCACCAA TCCACACAAA TTCAGCCACA 5040 ACATCACCCA ACACAAAGTC AGAAACACAC CACACAACAG CACAAACCAA AGGCAGAACC 5100 ACCACCTCAA CACAGACCAA CAAGCCGAGC ACAAAACCAC GCCTAAAAAA TCCACCAAAA 5160 AAACCAAAAG ATGATTACCA TTTTGAAGTG TTCAACTTCG TTCCCTGTAG TATATGTGGC 5220 AACAATCAAC TTTGCAAATC CATCTGTAAA ACAATACCAA GCAACAAACC AAAGAAGAAA 5280

CCAACCATCA AACCCACAAA CAAACCAACC ACCAAAACCA CAAACAAAAG AGACCCAAAA 5340 ACACCAGCCA AAACGACGAA AAAAGAAACT ACCACCAACC CAACAAAAAA ACCAACCCTC 5400 ACGACCACAG AAAGAGACAC CAGCACCTCA CAATCCACTG TGCTCGACAC AACCACATTA 5460 GAACACAAA TCCAACAGCA ATCCCTCCAC TCAACCACCC CCGAAAACAC ACCCAACTCC 5520 ACACAAACAC CCACAGCATC CGAGCCCTCT ACATCAAATT CCACCCAAAA TACCCAATCA 5580 CATGCTTAGT TATTCAAAAA CTACATCTTA GCAGAAAACC GTGACCTATC AAGCAAGAAC 5640 GAAATTAAAC CTGGGGCAAA TAACCATGGA GCTGCTGATC CACAGGTTAA GTGCAATCTT 5700 CCTAACTCTT GCTATTAATG CATTGTACCT CACCTCAAGT CAGAACATAA CTGAGGAGTT 5760 TTACCAATCG ACATGTAGTG CAGTTAGCAG AGGTTATTTT AGTGCTTTAA GAACAGGTTG 5820 GTATACCAGT GTCATAACAA TAGAATTAAG TAATATAAAA GAAACCAAAT GCAATGGAAC 5880 TGACACTAAA GTAAAACTTA TAAAACAAGA ATTAGATAAG TATAAGAATG CAGTGACAGA 5940 ATTACAGCTA CTTATGCAAA ACACACCAGC TGCCAACAAC CGGGCCAGAA GAGAAGCACC 6000 ACAGTATATG AACTATACAA TCAATACCAC TAAAAACCTA AATGTATCAA TAAGCAAGAA 6060 GAGGAAACGA AGATTTCTGG GCTTCTTGTT AGGTGTAGGA TCTGCAATAG CAAGTGGTAT 6120 AGCTGTATCC AAAGTTCTAC ACCTTGAAGG AGAAGTGAAC AAGATCAAAA ATGCTTTGTT 6180 ATCTACAAAC AAAGCTGTAG TCAGTCTATC AAATGGGGTC AGTGTTTTAA CCAGCAAAGT 6240 GTTAGATCTC AAGAATTACA TAAATAACCA ATTATTACCC ATAGTAAATC AACAGAGCTG 6300 TCGCATCTCC AACATTGAAA CAGTTATAGA ATTCCAGCAG AAGAACAGCA GATTGTTGGA 6360 AATCAACAGA GAATTCAGTG TCAATGCAGG TGTAACAACA CCTTTAAGCA CTTACATGTT 6420 AACAAACAGT GAGTTACTAT CATTGATCAA TGATATGCCT ATAACAAATG ATCAGAAAAA 6480 ATTAATGTCA AGCAATGTTC AGATAGTAAG GCAACAAAGT TATTCTATCA TGTCTATAAT 6540 AAAGGAAGAA GTCCTTGCAT ATGTTGTACA GCTACCTATC TATGGTGTAA TAGATACACC 6600 TTGCTGGAAA TTACACACAT CACCTCTATG CACCACCAAC ATCAAAGAAG GATCAAATAT 6660 TTGTTTAACA AGGACTGATA GAGGATGGTA TTGTGATAAT GCAGGATCAG TATCCTTCTT 6720 TCCACAGGCT GACACTTGTA AAGTACAGTC CAATCGAGTA TTTTGTGACA CTATGAACAG 6780 TTTGACATTA CCAAGTGAAG TCAGCCTTTG TAACACTGAC ATATTCAATT CCAAGTATGA 6840 CTGCAAAATT ATGACATCAA AAACAGACAT AAGCAGCTCA GTAATTACTT CTCTTGGAGC 6900 TATAGTGTCA TGCTATGGTA AAACTAAATG CACTGCATCC AACAAAAATC GTGGGATTAT 6960 AAAGACATTT TCTAATGGTT GTGACTATGT GTCAAACAAA GGAGTAGATA CTGTGTCAGT 7020 GGGCAACACT TTATACTATG TAAACAAGCT GGAAGGCAAG AACCTTTATG TAAAAGGGGA 7080 ACCTATAATA AATTACTATG ACCCTCTAGT GTTTCCTTCT GATGAGTTTG ATGCATCAAT 7140 ATCTCAAGTC AATGAAAAAA TCAATCAAAG TTTAGCTTTT ATTCGTAGAT CTGATGAATT 7200 ACTACATAAT GTAAATACTG GCAAATCTAC TACAAATATT ATGATAACTA CAATTATTAT 7260 AGTAATCATT GTAGTATTGT TATCATTAAT AGCTATTGGT TTGCTGTTGT ATTGCAAAGC 7320 CAAAAACACA CCAGTTACAC TAAGCAAAGA CCAACTAAGT GGAATCAATA ATATTGCATT 7380

| CAGCAAATAG | ACAAAAAACC | ACCTGATCAT | GTTTCAACAA | CAGTCTGCTG | ATCACCAATC | 7440 |
|------------|------------|------------|--------------------------|------------|------------|------|
| CCAAATCAAC | CCATAACAAA | CACTTCAACA | TCACAGTACA | GGCTGAATCA | TITCTTCACA | 7500 |
| TCATGCTACC | CACACAACTA | AGCTAGATCC | TTAACTCATA | GTTACATAAA | AACCTCAAGT | 7560 |
| ATCACAATCA | AACACTAAAT | CAACACATCA | TTCACAAAAT | TAACAGCTGG | GGCAAATATG | 7620 |
| TCGCGAAGAA | ATCCTTGTAA | ATTTGAGATT | AGAGGTCATT | GCTTGAATGG | TAGAAGATGT | 7680 |
| CACTACAGTC | ATAATTACTT | TGAATGGCCT | CCTCATGCCT | TACTAGTGAG | GCAAAACTTC | 7740 |
| ATGTTAAACA | AGATACTCAA | GTCAATGGAC | AAAAGCATAG | ACACTTTGTC | TGAAATAAGT | 7800 |
| GGAGCTGCTG | AACTGGACAG | AACAGAAGAA | TATGCTCTTG | GTATAGTTGG | AGTGCTAGAG | 7860 |
| AGTTACATAG | GATCTATAAA | CAACATAACA | AAACAATCAG | CATGTGTTGC | TATGAGTAAA | 7920 |
| CTTCTTATTG | AGATCAATAG | TGATGACATT | AAAAAGC _Ç TGA | GAGATAATGA | AGAACCCAAT | 7980 |
| TCACCTAAGA | TAAGAGTGTA | CAATACTGTT | ATATCATACA | TTGAGAGCAA | TAGAAAAAAC | 8040 |
| AACAAGCAAA | CAATCCATCT | GCTCAAAAGA | CTACCAGCAG | ACGTGCTGAA | GAAGACAATA | 8100 |
| AAAAACACAT | TAGATATCCA | CAAAAGCATA | ATCATAAGCA | ACCCAAAAGA | GTCAACCGTG | 8160 |
| AATGATCAAA | ATGACCAAAC | CAAAAATAAT | GATATTACCG | GATAAATATC | CTTGTAGTAT | 8220 |
| ATCATCCATA | TTGATTTCAA | GTGAAAGCAT | GATTGCTACA | TTCAATCATA | AAAACATATT | 8280 |
| ACAATTTAAC | CATAACCATT | TGGATAACCA | CCAGCGTTTA | ТТАААТААТА | TATTTGATGA | 8340 |
| AATTCATTGG | ACACCTAAAA | ACTTATTAGA | TGCCACTCAA | CAATTTCTCC | AACATCTTAA | 8400 |
| CATCCCTGAA | GATATATATA | CAATATATAT | ATTAGTGTCA | TAATGCTTGG | CCATAACGAT | 8460 |
| TCTATATCAT | CCAACCATAA | AACTATCTTA | ATAAGGTTAT | GGGACAAAAT | GGATCCCATT | 8520 |
| ATTAATGGAA | ACTCTGCTAA | TGTGTATCTA | ACTGATAGTT | ATTTAAAAGG | TGTTATCTCT | 8580 |
| TTTTCAGAAT | GTAATGCTTT | AGGGAGTTAC | CTTTTTAACG | GCCCTTATCT | CAAAAATGAT | 8640 |
| TACACCAACT | TAATTAGTAG | ACAAAGTCCA | CTACTAGAGC | ATATGAATCT | TAAAAAACTA | 8700 |
| ACTATAACAC | AGTCATTAAT | ATCTAGATAT | CATAAAGGTG | AACTGAAATT | AGAAGAACCA | 8760 |
| ACTTATTTCC | AGTCATTACT | TATGACATAT | AAAAGCATGT | CCTCGTCTGA | ACAAATTGCT | 8820 |
| ACAACTAACT | TACTTAAAAA | AATAATACGA | AGAGCTATAG | AAATAAGTGA | TGTAAAGGTG | 8880 |
| TACGCCATCT | TGAATAAACT | AGGACTAAAG | GAAAAGGACA | GAGTTAAGCC | CAACAATAAT | 8940 |
| TCAGGTGATG | AAAACTCAGT | ACTTACAACT | ATAATTAAAG | ATGATATACT | TTCGGCTGTG | 9000 |
| GAAAGCAATC | AATCATATAC | AAATTCAGAC | AAAAATCACT | CAGTAAATCA | AAATATCACT | 9060 |
| ATCAAAACAA | CACTCTTGAA | AAAATTGATG | TGTTCAATGC | AACATCCTCC | ATCATGGTTA | 9120 |
| ATACACTGGT | TCAATTTATA | TACAAAATTA | AATAACATAT | TAACACAATA | TCGATCAAAT | 9180 |
| GAGGTAAAAA | GTCATGGGTT | TATATTAATA | GATAATCAAA | CTTTAAGTGG | TTTTCAGTTT | 9240 |
| ATTTTAAATC | AATATGGTTG | TATCGTTTAT | CATAAAGGAC | TCAAAAAAA | CACAACTACT | 9300 |
| ACTTACAATC | AATTTTTAAC | ATGGAAAGAC | ATCAGCCTTA | GCAGATTAAA | TGTTTGCTTA | 9360 |
| ATTACTTGGA | TAAGTAATTG | TTTGAATACA | TTAAATAAAA | GCTTAGGGCT | GAGATGTGGA | 9420 |
| TTCAATAATG | TTGTGTTATC | ACAATTATTT | CTTTATGGAG | ATTGTATACT | GAAATTATTT | 9480 |

| CATAATGAAG GCTTCTACAT | AATAAAAGAA | GTAGAGGGAT | TTATTATGTC | TTTAATTCTA | 9540 |
|-----------------------|------------|------------|------------|------------|-------|
| AACATAACAG AAGAAGATCA | ATTTAGGAAA | CGATTTTATA | ATAGCATGCT | AAATAACATC | 9600 |
| ACAGATGCAG CTATTAAGGĆ | TCAAAAGAAC | СТАСТАТСАА | GGGTATGTCA | CACTITATTA | 9660 |
| GACAAGACAG TGTCTGATAA | TATCATAAAT | GGTAAATGGA | TAATCCTATT | AAGTAAATTT | 9720 |
| CTTAAATTGA TTAAGCTTGC | AGGTGATAAT | AATCTCAATA | ATTTGAGTGA | GCTATATTTT | 9780 |
| CTCTTCAGAA TCTTTGGACA | TCCAATGGTT | GATGAAAGAC | AAGCAATGGA | TGCTGTAAGA | 9840 |
| ATTAACTGTA ATGAAACTAA | GTTCTACTTA | TTAAGTAGTC | TAAGTACGTT | AAGAGGTGCT | 9900 |
| TTCATTTATA GAATCATAAA | AGGGTTTGTA | AATACCTACA | ACAGATGGCC | CACTTTAAGG | 9960 |
| AATGCTATTG TCCTACCTCT | AAGATGGTTA | AACTATTATA | AACTTAATAC | TTATCCATCT | 10020 |
| CTACTTGAAA TCACAGAAAA | TGATTTGATT | ATTTTATCAG | GATTGCGGTT | CTATCGTGAA | 10080 |
| TTTCATCTGC CTAAAAAAGT | GGATCTTGAA | ATGATAATAA | ATGACAAAGC | CATTTCACCT | 10140 |
| CCAAAAGATC TAATATGGAC | TAGTTTTCCT | AGAAATTACA | TGCCATCACA | TATACAAAAT | 10200 |
| TATATAGAAC ATGAAAAGTT | GAAGTTCTCT | GAAAGCGACA | GATCAAGAAG | AGTACTAGAG | 10260 |
| TATTACTTGA GAGATAATAA | ATTCAATGAA | TGCGATCTAT | ACAATTGTGT | AGTCAATCAA | 10320 |
| AGCTATCTCA ACAACTCTAA | TCACGTGGTA | TCACTAACTG | GTAAAGAAAG | AGAGCTCAGT | 10380 |
| GTAGGTAGAA TGTTTGCTAT | GCAACCAGGT | ATGTTTAGGC | AAATCCAAAT | CTTAGCAGAG | 10440 |
| AAAATGATAG CCGAAAATAT | TTTACAATTC | TTCCCTGAGA | GTTTGACAAG | ATATGGTGAT | 10500 |
| CTAGAGCTTC AAAAGATATT | AGAATTAAAA | GCAGGAATAA | GCAACAAGTC | AAATCGTTAT | 10560 |
| AATGATAACT ACAACAATTA | TATCAGTAAA | TGTTCTATCA | TTACAGATCT | TAGCAAATTC | 10620 |
| AATCAAGCAT TTAGATATGA | AACATCATGT | ATCTGCAGTG | ATGTATTAGA | TGAACTGCAT | 10680 |
| GGAGTACAAT CTCTGTTCTC | TTGGTTGCAT | TTAACAATAC | CTCTTGTCAC | AATAATATGT | 10740 |
| ACATATAGAC ATGCACCTCC | TTTCATAAAG | GATCATGTTG | TTAATCTTAA | TGAAGTTGAT | 10800 |
| GAACAAAGTG GATTATACAG | ATATCATATG | GGTGGTATTG | AGGGCTGGTG | TCAAAAACTG | 10860 |
| TGGACCATTG AAGCTATATC | ATTATTAGAT | CTAATATCTC | TCAAAGGGAA | ATTCTCTATC | 10920 |
| ACAGCTCTGA TAAATGGTGA | TAATCAGTCA | ATTGATATAA | GTAAACCAGT | TAGACTTATA | 10980 |
| GAGGGTCAGA CCCATGCTCA | AGCAGATTAT | TTGTTAGCAT | TAAATAGCCT | TAAATTGCTA | 11040 |
| TATAAAGAGT ATGCAGGTAT | AGGCCATAAG | CTTAAGGGAA | CAGAGACCTA | TATATCCCGA | 11100 |
| GATATGCAGT TCATGAGCAA | AACAATCCAG | CACAATGGAG | TGTACTATCC | AGCCAGTATC | 11160 |
| AAAAAAGTCC TGAGAGTAGG | TCCATGGATA | AATACAATAC | TTGATGATTT | TAAAGTTAGT | 11220 |
| TTAGAATCTA TAGGTAGCTT | AACACAGGAG | TTAGAATACA | GAGGGGAAAG | CTTATTATGC | 11280 |
| AGTTTAATAT TTAGGAACAT | TTGGTTATAC | AATCAAATTG | CTTTGCAACT | CCGAAATCAT | 11340 |
| GCATTATGTA ACAATAAGCT | ATATTTAGAT | ATATTGAAAG | TATTAAAACA | CTTAAAAACT | 11400 |
| TTTTTTAATC TTGATAGTAT | CGATATGGCG | TTATCATTGT | ATATGAATTT | GCCTATGCTG | 11460 |
| TTTGGTGGTG GTGATCCTAA | TTTGTTATAT | CGAAGCTTTT | ATAGGAGAAC | TCCAGACTTC | 11520 |
| CTTACAGAAG CTATAGTACA | TTCAGTGTTT | GTGTTGAGCT | ATTATACTGG | TCACGATTTA | 11580 |

CAAGATAAGC TCCAGGATCT TCCAGATGAT AGACTGAACA AATTCTTGAC ATGTGTCATC 11640 ACATTCGATA AAAATCCCAA TGCCGAGTTT GTAACATTGA TGAGGGATCC ACAGGCGTTA 11700 GGGTCTGAAA GGCAAGCTAA AATTACTAGT GAGATTAATA GATTAGCAGT AACAGAAGTC 11760 TTAAGTATAG CTCCAAACAA AATATTTTCT AAAAGTGCAC AACATTATAC TACCACTGAG 11820 ATTGATCTAA ATGACATTAT GCAAAATATA GAACCAACTT ACCCTCATGG ATTAAGAGTT 11880 GTTTATGAAA GTCTACCTTT TTATAAAGCA GAAAAAATAG TTAATCTTAT ATCAGGAACA 11940 AAATCCATAA CTAATATACT TGAAAAAACA TCAGCAATAG ATACAACTGA TATTAATAGG 12000 GCTACTGATA TGATGAGGAA AAATATAACT TTACTTATAA GGATACTTCC ACTAGATTGT 12060 AACAAAGACA AAAGAGAGTT ATTAAGTTTA GAAAATCTTA GTATAACTGA ATTAAGCAAG 12120 TATGTAAGAG AAAGATCTTG GTCATTATCC AATATAGTAG GAGTAACATC GCCAAGTATT 12180 ATGTTCACAA TGGACATTAA ATATACAACT AGCACTATAG CCAGTGGTAT AATTATAGAA 12240 AAATATAATG TTAATAGTTT AACTCGTGGT GAAAGAGGGC CTACTAAGCC ATGGGTAGGT 12300 TCATCTACGC AGGAGAAAA AACAATGCCA GTGTACAATA GACAAGTTTT AACCAAAAAG 12360 CAAAGAGACC AAATAGATTT ATTAGCAAAA TTAGACTGGG TATATGCATC CATAGACAAC 12420 AAAGATGAAT TCATGGAAGA ACTGAGTACT GGAACACTTG GACTGTCATA TGAAAAAGCC 12480 AAAAAGTTGT TTCCACAATA TCTAAGTGTC AATTATTTAC ACCGTTTAAC AGTCAGTAGT 12540 AGACCATGTG AATTCCCTGC ATCAATACCA GCTTATAGAA CAACAAATTA TCATTTCGAT 12600 ACTAGTCCTA TCAATCATGT ATTAACAGAA AAGTATGGAG ATGAAGATAT CGACATTGTG 12660 TTTCAAAATT GCATAAGTTT TGGTCTTAGC CTGATGTCGG TTGTGGAACA ATTCACAAAC 12720 ATATGTCCTA ATAGAATTAT TCTCATACCG AAGCTGAATG AGATACATTT GATGAAACCT 12780 CCTATATITA CAGGAGATGT TGATATCATC AAGTTGAAGC AAGTGATACA AAAACAGCAT 12840 ATGTTCCTAC CAGATAAAAT AAGTTTAACC CAATATGTAG AATTATTCCT AAGTAACAAA 12900 GCACTTAAAT CTGGATCTAA CATCAATTCT AATTTAATAT TAGTACATAA AATGTCTGAT 12960 13020 TATTTCATA ATGCTTATAT TTTAAGTACT AATTTAGCTG GACATTGGAT TCTAATTATT CAACTTATGA AAGATTCAAA AGGTATTTTT GAAAAAGATT GGGGAGAGGG GTACATAACT 13080 GATCATATGT TCATTAATTT GAATGTTTTC TTTAATGCTT ATAAGACTTA TTTGCTATGT 13140 TTTCATAAAG GTTATGGTAA AGCAAAATTA GAATGTGATA TGAACACTTC AGATCTTCTT 13200 TGTGTTTTGG AGTTAATAGA CAGTAGCTAC TGGAAATCTA TGTCTAAAGT TTTCCTAGAA 13260 CAAAAAGTCA TAAAATACAT AGTCAATCAA GACACAAGTT TGCATAGAAT AAAAGGCTGT 13320 CACAGTITTA AGTIGIGGIT TITAAAACGC CITAATAATG CIAAATTTAC CGTATGCCCT 13380 TGGGTTGTTA ACATAGATTA TCACCCAACA CATATGAAAG CTATATTATC TTACATAGAT 13440 13500 TTAGTTAGAA TGGGGTTAAT AAATGTAGAT AAATTAACCA TTAAAAATAA AAACAAATTC AATGATGAAT TTTACACATC AAATCTCTTT TACATTAGTT ATAACTTTTC AGACAACACT 13560 CATTTGCTAA CAAAACAAAT AAGAATTGCT AATTCAGAAT TAGAAGATAA TTATAACAAA 13620 CTATATCACC CAACCCCAGA AACTTTAGAA AATATATCAT TAATTCCTGT TAAAAGTAAT 13680

| AATAGTAACA | ÄACCTAAATT | TTGTATAAGT | GGAAATACCG | AATCTATAAT | GATGTCAACA | 13740 |
|------------|------------|------------|------------|------------|------------|-------|
| ТТСТСТААТА | AAATGCATAT | TAAATCTTCC | ACTGTTACCA | CAAGATTCAA | TTATAGCAAA | 13800 |
| CAAGACTTGT | ACAATTTATT | TCCAAATGTT | GTGATAGACA | GGATTATAGA | TCATTCAGGT | 13860 |
| AATACAGCAA | AATCTAACCA | ACTITACATC | ACCACTTCAC | ATCAGACATC | TTTAGTAAGG | 13920 |
| AATAGTGCAT | CACTTTATTG | CATGCTTCCT | TGGCATCATG | TCAATAGATT | TAACTTTGTA | 13980 |
| TTTAGTTCCA | CAGGATGCAA | GATCAGTATA | GAGTATATTT | TAAAAGATCT | TAAGATTAAG | 14040 |
| GACCCCAGTT | GTATAGCATT | CATAGGTGAA | GGAGCTGGTA | ACTTATTATT | ACGTACGGTA | 14100 |
| GTAGAACTTC | ATCCAGACAT | AAGATACATT | TACAGAAGTT | TAAAAGATTG | CAATGATCAT | 14160 |
| AGTTTACCTA | TTGAATTTCT | AAGATTATAC | AACGGGCATA | TAAACATAGA | TTATGGTGAG | 14220 |
| AATTTAACCA | TTCCTGCTAC | AGATGCAACT | AATAACATTC | ATTGGTCTTA | TTTACATATA | 14280 |
| AAATTTGCAG | AACCTATTAG | CATCTTTGTC | TGCGATGCTG | AATTACCTGT | TACAGCCAAT | 14340 |
| TGGAGTAAAA | TTATAATTGA | ATGGAGTAAG | CATGTAAGAA | AGTGCAAGTA | CTGTTCTTCT | 14400 |
| GTAAATAGAT | GCATTTTAAT | CGCAAAATAT | CATGCTCAAG | ATGATATTGA | TTTCAAATTA | 14460 |
| GATAACATTA | CTATATTAAA | AACTTACGTG | TGCCTAGGTA | GCAAGTTAAA | AGGATCTGAA | 14520 |
| GTTTACTTAG | TCCTTACAAT | AGGCCCTGCA | AATATACTTC | CTGTTTTTGA | TGTTGTGCAA | 14580 |
| AATGCTAAAT | TGATTTTTTC | AAGAACTAAA | AATTTCATTA | TGCCTAAAAA | AACTGACAAG | 14640 |
| GAATCTATCG | ATGCAAATAT | TAAAAGCTTA | ATACCTTTCC | TTTGTTACCC | TATAACAAAA | 14700 |
| AAAGGAATTA | AGACTTCATT | GTCAAAATTG | AAGAGTGTAG | TTAATGGGGA | TATATTATCA | 14760 |
| TATTCTATAG | CTGGACGTAA | TGAAGTATTC | AGCAACAAGC | TTATAAACCA | CAAGCATATG | 14820 |
| AATATCCTAA | AATGGCTAGA | TCATGTTTTA | AATTTTAGAT | CAGCTGAACT | TAATTACAAT | 14880 |
| CATTTATACA | TGATAGAGTC | CACATATCCT | TACTTAAGTG | AATTGTTAAA | TAGTTTAACA | 14940 |
| ACCAATGAGC | TCAAGAAACT | GATTAAAATA | ACAGGTAGTG | TACTATACAA | CCTTCCCAAC | 15000 |
| GAACAGTAAC | TTAAAATATC | ATTAACAAGT | TTGGTCAAAT | TTAGATGCTA | ACACATCATT | 15060 |
| ATATTATAGT | TATTAAAAAA | TATGCAAACT | TTTCAATAAT | TTAGCTTACT | GATTCCAAAA | 15120 |
| TTATCATTTT | ATTTTTAAGG | GGTTGAATAA | AAGTCTAAAA | CTAACAATGA | TACATGTGCA | 15180 |
| TTTACAACAC | AACGAGACAT | TAGTTTTTGA | CACTTTTTTT | CTCGT | | 15225 |

(2) INFORMATION FOR SEQ ID NO:3:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 33 base pairs
 (B) TYPE: nucleic acid
 (C) STRANDEDNESS: single
 (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: cDNA
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

(2) INFORMATION FOR SEQ ID NO:4:

| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 31 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear | |
|--|----|
| (ii) MOLECULE TYPE: cDNA | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4: | 21 |
| CCCGGGATAT TTTTTATTAA CTTATTTGAG T | 31 |
| (2) INFORMATION FOR SEQ ID NO:5: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear | |
| (ii) MOLECULE TYPE: cDNA | |
| | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5: | |
| GAAAGTATAT ATTATGTT | 18 |
| (2) INFORMATION FOR SEQ ID NO:6: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear | |
| (ii) MOLECULE TYPE: cDNA | |
| | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:6: | |
| TATATAAGCA CGATGATATG | 20 |
| (2) INFORMATION FOR SEQ ID NO:7: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 16 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear | |
| (ii) MOLECULE TYPE: cDNA | |
| | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:7: | |
| ACTCAAATAA GTTAAT | 16 |

| (2) | INFOR | MATION FOR SEQ ID NO:8: | |
|-----|--------|--|----|
| | (i) | SEQUENCE CHARACTERISTICS: (A) LENGTH: 14 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear | |
| | (ii) | MOLECULE TYPE: cDNA | |
| | (xi) | SEQUENCE DESCRIPTION: SEQ ID NO:8: | |
| TAA | TTATT | T GAGT | 14 |
| (2) | INFOR | MATION FOR SEQ ID NO:9: | |
| | (i) | SEQUENCE CHARACTERISTICS: (A) LENGTH: 28 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear | |
| | (ii) | MOLECULE TYPE: cDNA | |
| | | • | |
| | (xi) | SEQUENCE DESCRIPTION: SEQ ID NO:9: | |
| GAC | ACAACO | C ACAATGATAA TACACCAC | 28 |
| (2) | INFOR | MATION FOR SEQ ID NO:10: | |
| | (i) | SEQUENCE CHARACTERISTICS: (A) LENGTH: 32 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear | |
| | (ii) | MOLECULE TYPE: cDNA | |
| | | | |
| | (xi) | SEQUENCE DESCRIPTION: SEQ ID NO:10: | |
| CAT | CTCTAA | C CAAGGGAGTT AAATTTAAGT GG | 32 |
| (2) | INFOR | MATION FOR SEQ ID NO:11: | |
| | (i) | SEQUENCE CHARACTERISTICS: (A) LENGTH: 27 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear | |
| | (ii) | MOLECULE TYPE: cDNA | |
| | | | |
| | (xi) | SEQUENCE DESCRIPTION: SEQ ID NO:11: | |
| TTA | AGGAGA | AG ATATAAGATA GAAGATG | 27 |

| (2) INFORMATION FOR SEQ ID NO:12: | |
|--|----|
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 27 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear | |
| (ii) MOLECULE TYPE: cDNA | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:12: | |
| GTTTTATATT AACTAATGGT GTTAGTG | 27 |
| (2) INFORMATION FOR SEQ ID NO:13: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 33 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear | |
| (ii) MOLECULE TYPE: cDNA | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:13: | |
| TTATAATTGC AGCCATCATA TTCATAGCCT CGG | 33 |
| (2) INFORMATION FOR SEQ ID NO:14: | |
| (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 30 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear | |
| (ii) MOLECULE TYPE: cDNA | |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:14: | |
| GTGAAGTTGA GATTACAATT GCCAGAATGG | 30 |